SEQUENCE LISTING

<110> Japan EnviroChemicals, Ltd.

<120> A protein binding to plasticizers

<130> 09622

<150> JP 2003-110877
<151> 2003-4-15

<160> 27

<170> PatentIn version 3.1

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Glu Val His Leu Val Glu Ser Gly Gly Asp Leu Val Arg Pro Gly Gly
1               5                  10                  15

tcc ctg aaa ctc tcc tgt gca gcc tct gga ttc act ttc gga agt tat       96
Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Gly Ser Tyr
            20                  25                  30

ggc atg tct tgg gtt cgc cag act gca gac aag agg ctg gag tgg gtc      144
Gly Met Ser Trp Val Arg Gln Thr Ala Asp Lys Arg Leu Glu Trp Val
        35                  40                  45

gca acc att tat agt ggt ggt ttt tac acc tac tat cca gac agt gtg      192
Ala Thr Ile Tyr Ser Gly Gly Phe Tyr Thr Tyr Tyr Pro Asp Ser Val
    50                  55                  60

agg gga cga ttc acc atc tcc aga gac aat gtc aag gaa atc gtg tat      240
Arg Gly Arg Phe Thr Ile Ser Arg Asp Asn Val Lys Glu Ile Val Tyr
65                  70                  75                  80

ctg caa atg agc agt ctg aag tct gag gac aca gcc atg tat tac tgt      288
Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met Tyr Tyr Cys
                85                  90                  95

gca aga cgg acg gta gta tct acg gac tat act ttg gac tac tgg ggt      336
Ala Arg Arg Thr Val Val Ser Thr Asp Tyr Thr Leu Asp Tyr Trp Gly
            100                 105                 110

caa gga acc tca gtc atc gtc tcc tca                                 363
Gln Gly Thr Ser Val Ile Val Ser Ser
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Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Gly Ser Tyr
            20                  25                  30

Gly Met Ser Trp Val Arg Gln Thr Ala Asp Lys Arg Leu Glu Trp Val
        35                  40                  45

Ala Thr Ile Tyr Ser Gly Gly Phe Tyr Thr Tyr Tyr Pro Asp Ser Val
    50                  55                  60
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Arg Gly Arg Phe Thr Ile Ser Arg Asp Asn Val Lys Glu Ile Val Tyr
65                  70                  75                  80

Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met Tyr Tyr Cys
                85                  90                  95

Ala Arg Arg Thr Val Val Ser Thr Asp Tyr Thr Leu Asp Tyr Trp Gly
            100                 105                 110

Gln Gly Thr Ser Val Ile Val Ser Ser
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1               5                   10                  15

gac aga gtc acc atc agt tgc cgg cca agt cag gac atc agc aat ttt      96
Asp Arg Val Thr Ile Ser Cys Arg Pro Ser Gln Asp Ile Ser Asn Phe
            20                  25                  30

tta aac tgg ttt cag cag aaa cca gat gga act gtt gaa gtc ctg atc     144
Leu Asn Trp Phe Gln Gln Lys Pro Asp Gly Thr Val Glu Val Leu Ile
        35                  40                  45

tgc tac aca tta aga atg cac tta gga gtc cca tca acg ttc agt ggc     192
Cys Tyr Thr Leu Arg Met His Leu Gly Val Pro Ser Thr Phe Ser Gly
    50                  55                  60

tgt gtg tct gga aca tat tat act ctc acc agt agc aac ctg gaa caa     240
Cys Val Ser Gly Thr Tyr Tyr Thr Leu Thr Ser Ser Asn Leu Glu Gln
65                  70                  75                  80

gaa gat ata gac act tcc ttt gcc att agg att ata cgc gtg ctc acg     288
Glu Asp Ile Asp Thr Ser Phe Ala Ile Arg Ile Ile Arg Val Leu Thr
                85                  90                  95

gtc ggt gca ggg acc acg ctg gag ctg aaa                             318
Val Gly Ala Gly Thr Thr Leu Glu Leu Lys
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Asp Arg Val Thr Ile Ser Cys Arg Pro Ser Gln Asp Ile Ser Asn Phe
            20                  25                  30

Leu Asn Trp Phe Gln Gln Lys Pro Asp Gly Thr Val Glu Val Leu Ile
        35                  40                  45

Cys Tyr Thr Leu Arg Met His Leu Gly Val Pro Ser Thr Phe Ser Gly
    50                  55                  60

Cys Val Ser Gly Thr Tyr Tyr Thr Leu Thr Ser Ser Asn Leu Glu Gln
65                  70                  75                  80

Glu Asp Ile Asp Thr Ser Phe Ala Ile Arg Ile Ile Arg Val Leu Thr
                85                  90                  95

Val Gly Ala Gly Thr Thr Leu Glu Leu Lys
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<210> 6
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<212> PRT
<213> Artificial

<220>
<223> Linker

<400> 6
Gly Ser Thr Ser Gly Ser Gly Lys Ser Ser Glu Gly Lys Gly
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<211> 18
<212> PRT
<213> Artificial

<220>
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<400> 7
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Lys Gly

<210> 8
<211> 12
<212> PRT
<213> Artificial

<220>
<223> Linker

<400> 8
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1 5 10

<210> 9
<211> 18
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<213> Artificial

<220>
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<400> 9
Gly Ser Thr Ser Gly Ser Gly Lys Pro Gly Ser Gly Glu Gly Ser Thr
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Lys Gly

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<220>
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<400> 11
acactgctgg acagggat 18

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<220>
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<400> 12
ggatcccggg agtacccctt gaccaggc 28

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<212> DNA
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<220>
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<400> 13
gttgaagctc ttgacaat 18

<210> 14
<211> 27
<212> DNA
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<220>
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<400> 14
ggatcccggg tggatggtgg gaagatg 27

<210> 15
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<212> DNA
<213> Artificial

<220>
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<400> 15
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<212> DNA
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<220>
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<400> 16
ggccacgcgt cgactagtac 20

<210> 17
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<220>
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<400> 17
actagtcgac atggtrtccw casctcagtt ccttg 35

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ggaaacagct atgaccatg 19

<210> 19
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<400> 19
gtaaaacgac ggccagt 17

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<213> Artificial

<220>
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<400> 20
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<210> 21
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<400> 21
ccgccggatc cacctccgcc tgaaccgcct ccacctgagg agacgatgac tgaggttcc 59

<210> 22

<211> 56
<212> DNA
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<220>
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<400> 22
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<210> 23
<211> 67
<212> DNA
<213> Artificial

<220>
<223> Primer

<400> 23
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tccctgc 67

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tct ctg tct ctc acc tgt tct gtc act ggc tac tcc atc acc agt ggt 96
Ser Leu Ser Leu Thr Cys Ser Val Thr Gly Tyr Ser Ile Thr Ser Gly
20 25 30

tat tac tgg aat tgg atc cgg caa ttt cca gga aac aaa ctg gat tgg 144
Tyr Tyr Trp Asn Trp Ile Arg Gln Phe Pro Gly Asn Lys Leu Asp Trp
35 40 45

atg ggc cat ata agt tac gac ggt aac aat aac tac aac cca tct ctc 192
Met Gly His Ile Ser Tyr Asp Gly Asn Asn Asn Tyr Asn Pro Ser Leu
50 55 60

aaa aat cga atc tcc atc act cgt gac aca tct aag aac cag ttt ttc 240
Lys Asn Arg Ile Ser Ile Thr Arg Asp Thr Ser Lys Asn Gln Phe Phe
65 70 75 80

ctg aag ttg aat tct gtg act act gag gac aca gat aca tat tac tgt 288
Leu Lys Leu Asn Ser Val Thr Thr Glu Asp Thr Asp Thr Tyr Tyr Cys
85 90 95

tct atg atc ctc tat ggt atg gac tac tgg ggt cag gga acc tca gtc 336
Ser Met Ile Leu Tyr Gly Met Asp Tyr Trp Gly Gln Gly Thr Ser Val
100 105 110

acc gtc tcc tca 348
Thr Val Ser Ser
115

<210> 25
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<212> PRT
<213> Mus musculus

<400> 25
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Ser Leu Ser Leu Thr Cys Ser Val Thr Gly Tyr Ser Ile Thr Ser Gly
20 25 30

Tyr Tyr Trp Asn Trp Ile Arg Gln Phe Pro Gly Asn Lys Leu Asp Trp
35 40 45

Met Gly His Ile Ser Tyr Asp Gly Asn Asn Asn Tyr Asn Pro Ser Leu
50 55 60

Lys Asn Arg Ile Ser Ile Thr Arg Asp Thr Ser Lys Asn Gln Phe Phe
65 70 75 80

Leu Lys Leu Asn Ser Val Thr Thr Glu Asp Thr Asp Thr Tyr Tyr Cys
85 90 95

Ser Met Ile Leu Tyr Gly Met Asp Tyr Trp Gly Gln Gly Thr Ser Val
100 105 110

Thr Val Ser Ser
115

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<212> DNA
<213> Mus musculus

<220>
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<222> (1)..(324)

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Gln Ile Val Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Leu Gly
1 5 10 15

gaa cgg gtc acc atg acc tgc act gcc agc tca agt gta agt tcc agt 96
Glu Arg Val Thr Met Thr Cys Thr Ala Ser Ser Ser Val Ser Ser Ser
20 25 30

tac ttg cac tgg tac cag cag aag cca gga tcc tcc ccc aaa ctc tgc 144
Tyr Leu His Trp Tyr Gln Gln Lys Pro Gly Ser Ser Pro Lys Leu Cys
35 40 45

att tat agc aca tcc aac ctg gct tct gga gtc cca act cgc ttc agt 192
Ile Tyr Ser Thr Ser Asn Leu Ala Ser Gly Val Pro Thr Arg Phe Ser
50 55 60

ggc agt ggg tct ggg acc tct tac tct ctc aca ata agc agc atg gag 240
Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu
65 70 75 80

gct gaa gat gct gcc act tat tac tgc cac cag tat cat cgt tcc cca 288
Ala Glu Asp Ala Ala Thr Tyr Tyr Cys His Gln Tyr His Arg Ser Pro
85 90 95

ccc acg ttc ggc tcg ggg aca aag ttg gaa ata aaa 324
Pro Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys
100 105

<210> 27
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<212> PRT
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<400> 27

Gln Ile Val Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Leu Gly
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20 25 30

Tyr Leu His Trp Tyr Gln Gln Lys Pro Gly Ser Ser Pro Lys Leu Cys
35 40 45

Ile Tyr Ser Thr Ser Asn Leu Ala Ser Gly Val Pro Thr Arg Phe Ser
50 55 60

Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu
65 70 75 80

Ala Glu Asp Ala Ala Thr Tyr Tyr Cys His Gln Tyr His Arg Ser Pro
85 90 95

Pro Thr Phe Gly Ser Gly Thr Lys Leu Glu Ile Lys
100 105

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<212> DNA
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<220>
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<400> 28
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<210> 29
<211> 40
<212> DNA
<213> Artificial

<220>
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<400> 29
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<210> 30
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<212> DNA
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<211> 61
<212> DNA
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<210> 32
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<212> DNA
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gctcaacttt cttgtcgact ttatcatcat catctttata atcttttatt tccaactttg 60

tccccg 66

<210> 34
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<213> Artificial

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<400> 34
Gly Gly Gly Gly Ser
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